

# Accessing Data within Pandas

## Introduction

In this lesson, we're going to dig into various methods for accessing data from our Pandas Series and DataFrames.

## Objectives

You will be able to:

- Use pandas methods and attributes to access information about a dataset
- Index pandas dataframes with `.loc`, `.iloc`, and column names
- Use a boolean mask to index pandas series and dataframes

## Importing pandas and the data

First, let's make sure we import `pandas` as `pd`.

```
In [1]: import pandas as pd
```

To show how to access data with Pandas, let's use the `wine` dataset in the scikit-learn library. Don't worry about the code below. We're essentially just making sure you have access to the `wine` dataset.

The data contained in the wine dataset are the results of a chemical analysis of wines grown in Italy. It contains the quantities of 13 wine constituents.

```
In [2]: from sklearn.datasets import load_wine

data = load_wine()
df = pd.DataFrame(data.data, columns=data.feature_names)
```

Great! Our data set is now stored in the variable `df`. As you know, you can look at its elements by using `df` or `print(df)`.

```
In [3]: print(df)
```

	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	total_pheno
0	14.23	1.71	2.43	15.6	127.0	2.
1	13.20	1.78	2.14	11.2	100.0	2.
2	13.16	2.36	2.67	18.6	101.0	2.
3	14.37	1.95	2.50	16.8	113.0	3.
4	13.24	2.59	2.87	21.0	118.0	2.
5	14.20	1.76	2.45	15.2	112.0	3.
6	14.39	1.87	2.45	14.6	96.0	2.
7	14.06	2.15	2.61	17.6	121.0	2.
8	14.83	1.64	2.17	14.0	97.0	2.

Now, what if you want to see only a few lines of the data, based on certain constraints? You'll learn how to access data in this lesson!

## Methods and attributes to access data information

It won't be a surprise that our `df` object is a Pandas DataFrame object. Let's verify this using the `type()` function:

```
In [4]: type(df)
```

```
Out[4]: pandas.core.frame.DataFrame
```

There are some methods and attributes associated with Pandas objects (both DataFrames *and* series!) which make retrieving information from the data particularly easy. Some commonly used methods:

- `.head()`
- `.tail()`

And attributes:

- `.index`
- `.columns`
- `.dtypes`
- `.shape`

## Some methods: `.head()`, `.tail()`, and `.info()`

By using `.head()` and `.tail()`, you can select the first  $n$  rows from your dataframe. The default  $n$  is 5, but you can change this value inside the parentheses. For example:

```
In [5]: # First 5 rows of df
df.head()
```

```
Out[5]:
```

	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	total_phenols	flavanoids	nonflavanoid_phenols
0	14.23	1.71	2.43	15.6	127.0	2.80	3.06	
1	13.20	1.78	2.14	11.2	100.0	2.65	2.76	
2	13.16	2.36	2.67	18.6	101.0	2.80	3.24	
3	14.37	1.95	2.50	16.8	113.0	3.85	3.49	
4	13.24	2.59	2.87	21.0	118.0	2.80	2.69	

```
In [6]: # last 3 rows of df
df.tail(3)
```

```
Out[6]:
```

	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	total_phenols	flavanoids	nonflavanoid_phenols
175	13.27	4.28	2.26	20.0	120.0	1.59	0.69	
176	13.17	2.59	2.37	20.0	120.0	1.65	0.68	
177	14.13	4.10	2.74	24.5	96.0	2.05	0.76	

To get a concise summary of the dataframe, you can use `.info()` :

```
In [7]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 178 entries, 0 to 177
Data columns (total 13 columns):
alcohol                178 non-null float64
malic_acid             178 non-null float64
ash                   178 non-null float64
alcalinity_of_ash      178 non-null float64
magnesium              178 non-null float64
total_phenols          178 non-null float64
flavanoids             178 non-null float64
nonflavanoid_phenols   178 non-null float64
proanthocyanins        178 non-null float64
color_intensity        178 non-null float64
hue                   178 non-null float64
od280/od315_of_diluted_wines 178 non-null float64
proline                178 non-null float64
dtypes: float64(13)
memory usage: 18.2 KB
```

## Some attributes

Using `.index`, you can access the index or row labels of the DataFrame.

```
In [8]: df.index
```

```
Out[8]: RangeIndex(start=0, stop=178, step=1)
```

Using `.columns`, you can access the column labels of the DataFrame.

```
In [9]: df.columns
```

```
Out[9]: Index(['alcohol', 'malic_acid', 'ash', 'alcalinity_of_ash', 'magnesium',  
              'total_phenols', 'flavanoids', 'nonflavanoid_phenols',  
              'proanthocyanins', 'color_intensity', 'hue',  
              'od280/od315_of_diluted_wines', 'proline'],  
             dtype='object')
```

Using `.dtypes` returns the data types of all columns in the DataFrame (compare with `.info()` !)

```
In [10]: df.dtypes
```

```
Out[10]: alcohol                float64  
malic_acid                    float64  
ash                          float64  
alcalinity_of_ash            float64  
magnesium                    float64  
total_phenols                float64  
flavanoids                   float64  
nonflavanoid_phenols         float64  
proanthocyanins              float64  
color_intensity              float64  
hue                          float64  
od280/od315_of_diluted_wines float64  
proline                      float64  
dtype: object
```

`.shape` returns a tuple representing the dimensionality (in `(rows, columns)`) of the DataFrame.

```
In [11]: df.shape
```

```
Out[11]: (178, 13)
```

## Selecting DataFrame information

In the previous section, we deliberately omitted 2 very important attributes:

- `.iloc` , which is a Pandas DataFrame indexer used for integer-location based indexing / selection by position
- `.loc` , which has two use cases:
  - Selecting by label / index
  - Selecting with a boolean / conditional lookup

## `.iloc`

You can use `.iloc` to select single rows. To select the 4th row, you can use `.iloc[3]` like:

```
In [12]: df.iloc[3]
```

```
Out[12]: alcohol      14.37
malic_acid      1.95
ash      2.50
alcalinity_of_ash      16.80
magnesium      113.00
total_phenols      3.85
flavanoids      3.49
nonflavanoid_phenols      0.24
proanthocyanins      2.18
color_intensity      7.80
hue      0.86
od280/od315_of_diluted_wines      3.45
proline      1480.00
Name: 3, dtype: float64
```

You can use a colon to select several rows. Note that you'll use a structure `.iloc[a:b]` where the row with index `a` will be included in the selection and the row with index `b` is excluded.

```
In [13]: df.iloc[5:8]
```

```
Out[13]:
```

	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	total_phenols	flavanoids	nonflavanoid_phenols
5	14.20	1.76	2.45	15.2	112.0	3.27	3.39	
6	14.39	1.87	2.45	14.6	96.0	2.50	2.52	
7	14.06	2.15	2.61	17.6	121.0	2.60	2.51	

Next, you can use `,` to perform *column* selections based on their index as well. The command below selects full columns 3-6:

```
In [14]: df.iloc[:, 3:7]
```

Out[14]:

	alcalinity_of_ash	magnesium	total_phenols	flavanoids
0	15.6	127.0	2.80	3.06
1	11.2	100.0	2.65	2.76
2	18.6	101.0	2.80	3.24
3	16.8	113.0	3.85	3.49
4	21.0	118.0	2.80	2.69
5	15.2	112.0	3.27	3.39
6	14.6	96.0	2.50	2.52
7	17.6	121.0	2.60	2.51
8	14.0	97.0	2.80	2.98
9	16.0	98.0	2.98	3.15
10	18.0	105.0	2.95	3.32
11	16.8	95.0	2.20	2.43
12	16.0	89.0	2.60	2.76
13	11.4	91.0	3.10	3.69
14	12.0	102.0	3.30	3.64
15	17.2	112.0	2.85	2.91
16	20.0	120.0	2.80	3.14
17	20.0	115.0	2.95	3.40
18	16.5	108.0	3.30	3.93
19	15.2	116.0	2.70	3.03
20	16.0	126.0	3.00	3.17
21	18.6	102.0	2.41	2.41
22	16.6	101.0	2.61	2.88
23	17.8	95.0	2.48	2.37
24	20.0	96.0	2.53	2.61
25	25.0	124.0	2.63	2.68
26	16.1	93.0	2.85	2.94
27	17.0	94.0	2.40	2.19
28	19.4	107.0	2.95	2.97
29	16.0	96.0	2.65	2.33
...	...	...	...	...
148	21.5	92.0	1.93	0.76
149	21.5	113.0	1.41	1.39
150	24.0	123.0	1.40	1.57
151	22.0	112.0	1.48	1.36

	alcalinity_of_ash	magnesium	total_phenols	flavanoids
152	25.5	116.0	2.20	1.28
153	18.5	98.0	1.80	0.83
154	20.0	103.0	1.48	0.58
155	22.0	93.0	1.74	0.63
156	19.5	89.0	1.80	0.83
157	27.0	97.0	1.90	0.58
158	25.0	98.0	2.80	1.31
159	22.5	89.0	2.60	1.10
160	21.0	88.0	2.30	0.92
161	20.0	107.0	1.83	0.56
162	22.0	106.0	1.65	0.60
163	18.5	106.0	1.39	0.70
164	22.0	90.0	1.35	0.68
165	22.5	88.0	1.28	0.47
166	23.0	111.0	1.70	0.92
167	19.5	88.0	1.48	0.66
168	24.5	105.0	1.55	0.84
169	25.0	112.0	1.98	0.96
170	19.0	96.0	1.25	0.49
171	19.5	86.0	1.39	0.51
172	20.0	91.0	1.68	0.70
173	20.5	95.0	1.68	0.61
174	23.0	102.0	1.80	0.75
175	20.0	120.0	1.59	0.69
176	20.0	120.0	1.65	0.68
177	24.5	96.0	2.05	0.76

178 rows × 4 columns

Last but not least, you can perform column and row selections at once:



```
In [15]: df.iloc[5:10, 3:9]
```

```
Out[15]:
```

	alkalinity_of_ash	magnesium	total_phenols	flavanoids	nonflavanoid_phenols	proanthocyanins
5	15.2	112.0	3.27	3.39	0.34	1.97
6	14.6	96.0	2.50	2.52	0.30	1.98
7	17.6	121.0	2.60	2.51	0.31	1.25
8	14.0	97.0	2.80	2.98	0.29	1.98
9	16.0	98.0	2.98	3.15	0.22	1.85

## **.loc**

### **a) .loc label-based indexing**

You can `.loc` to select columns based on their (row index and) column name. Examples:

```
In [16]: df.loc[:, 'magnesium']
```

```
Out[16]: 0      127.0
         1      100.0
         2      101.0
         3      113.0
         4      118.0
         5      112.0
         6       96.0
         7      121.0
         8       97.0
         9       98.0
        10      105.0
        11       95.0
        12       89.0
        13       91.0
        14      102.0
        15      112.0
        16      120.0
        17      115.0
        18      108.0
        19      116.0
        20      126.0
        21      102.0
        22      101.0
        23       95.0
        24       96.0
        25      124.0
        26       93.0
        27       94.0
        28      107.0
        29       96.0
        ...
       148       92.0
       149      113.0
       150      123.0
       151      112.0
       152      116.0
       153       98.0
       154      103.0
       155       93.0
       156       89.0
       157       97.0
       158       98.0
       159       89.0
       160       88.0
       161      107.0
       162      106.0
       163      106.0
       164       90.0
       165       88.0
       166      111.0
       167       88.0
       168      105.0
       169      112.0
       170       96.0
       171       86.0
       172       91.0
       173       95.0
```

```
174      102.0
175      120.0
176      120.0
177       96.0
Name: magnesium, Length: 178, dtype: float64
```

An alternative method here is simply calling `df['magnesium']` !

```
In [17]: df.loc[7:16, 'magnesium']
```

```
Out[17]: 7      121.0
          8       97.0
          9       98.0
         10      105.0
         11       95.0
         12       89.0
         13       91.0
         14      102.0
         15      112.0
         16      120.0
Name: magnesium, dtype: float64
```

### b) boolean indexing using `.loc`

Sometimes you'd like to select certain rows in your dataset based on the value for a certain variable. Imagine you'd like to create a new DataFrame that only contains the wines with an alcohol percentage below 12. This can be done as follows:

```
In [18]: df.loc[df['alcohol'] < 12]
```

```
Out[18]:
```

	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	total_phenols	flavanoids	nonflavanoid_
74	11.96	1.09	2.30	21.0	101.0	3.38	2.14	
75	11.66	1.88	1.92	16.0	97.0	1.61	1.57	
77	11.84	2.89	2.23	18.0	112.0	1.72	1.32	
84	11.84	0.89	2.58	18.0	94.0	2.20	2.21	
87	11.65	1.67	2.62	26.0	88.0	1.92	1.61	
88	11.64	2.06	2.46	21.6	84.0	1.95	1.69	
94	11.62	1.99	2.28	18.0	98.0	3.02	2.26	
96	11.81	2.12	2.74	21.5	134.0	1.60	0.99	
103	11.82	1.72	1.88	19.5	86.0	2.50	1.64	
109	11.61	1.35	2.70	20.0	94.0	2.74	2.92	
110	11.46	3.74	1.82	19.5	107.0	3.18	2.58	
112	11.76	2.68	2.92	20.0	103.0	1.75	2.03	
113	11.41	0.74	2.50	21.0	88.0	2.48	2.01	
115	11.03	1.51	2.20	21.5	85.0	2.46	2.17	
116	11.82	1.47	1.99	20.8	86.0	1.98	1.60	
120	11.45	2.40	2.42	20.0	96.0	2.90	2.79	
121	11.56	2.05	3.23	28.5	119.0	3.18	5.08	
124	11.87	4.31	2.39	21.0	82.0	2.86	3.03	
127	11.79	2.13	2.78	28.5	92.0	2.13	2.24	

You can verify that simply using `df[df['alcohol'] < 12]`, you can obtain the same result!

However, the `.loc` attribute is useful if you'd only want the color intensity for the wines with an alcohol percentage below 12. You can obtain the result as follows:

```
In [19]: df.loc[df['alcohol'] < 12, ['color_intensity']]
```

```
Out[19]:
```

	color_intensity
74	3.21
75	3.80
77	2.65
84	3.05
87	2.60
88	2.80
94	3.25
96	2.50
103	2.06
109	2.65
110	2.90
112	3.80
113	3.08
115	1.90
116	1.95
120	3.25
121	6.00
124	2.80
127	3.00

## Selectors for series

Until now we've only really discussed Pandas DataFrames. Most of these methods and selectors are also applicable to Pandas Series. See how you can convert a one-column DataFrame into a Pandas Series:

```
In [20]: # Let's save our color intensity dataframe into an object col_intensity
col_intensity = df['color_intensity']
```

```
In [21]: type(col_intensity)
```

```
Out[21]: pandas.core.series.Series
```

Note how `col_intensity` is now a Pandas *Series*.

Many of the commands discussed before are readily applicable to series:

```
In [22]: col_intensity[0:3]
```

```
Out[22]: 0    5.64  
         1    4.38  
         2    5.68  
         Name: color_intensity, dtype: float64
```

```
In [23]: # Or col_intensity.loc[col_intensity > 8]  
         col_intensity[col_intensity > 8]
```

```
Out[23]: 18    8.700000  
         49    8.900000  
         144   8.210000  
         148   8.420000  
         149   9.400000  
         150   8.600000  
         151  10.800000  
         153  10.520000  
         156   9.010000  
         158  13.000000  
         159  11.750000  
         164   9.580000  
         166  10.680000  
         167  10.260000  
         168   8.660000  
         169   8.500000  
         171   9.899999  
         172   9.700000  
         175  10.200000  
         176   9.300000  
         177   9.200000  
         Name: color_intensity, dtype: float64
```

## Changing and setting values in DataFrames and series

### Changing values

Imagine that for some reason, you're not interested in the color intensity values for color intensities above 10, and simply want to set all color intensities to 10 when they are bigger than 10. You can use a selector method and then assign it a new value, just like this:

```
In [24]: df.loc[df['color_intensity'] > 10, 'color_intensity'] = 10
```

### Creating new columns

Now imagine that we want to create a new column named, "shade" which has a value, "light" when the `color_intensity` is below 7, and, "dark" when the intensity is  $> 7$ . This can be done as follows:

```
In [25]: df.loc[df['color_intensity'] > 7, 'shade'] = 'dark'  
df.loc[df['color_intensity'] <= 7, 'shade'] = 'light'
```

If you now look at the output of `df.shape`, you will notice that `df` now has 14 columns.

```
In [26]: df.shape
```

```
Out[26]: (178, 14)
```

## Summary

We've introduced a range of techniques for accessing information in Pandas Series and DataFrames, selecting rows and columns, changing values, and creating new columns! Now, it's time for some practice! Let's start working on a lab where you will get a chance to practice some of these methods!